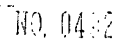


Figure 1

$\frac{d}{dt} \left(\frac{\partial L}{\partial \dot{x}} \right) = \frac{\partial L}{\partial x}$



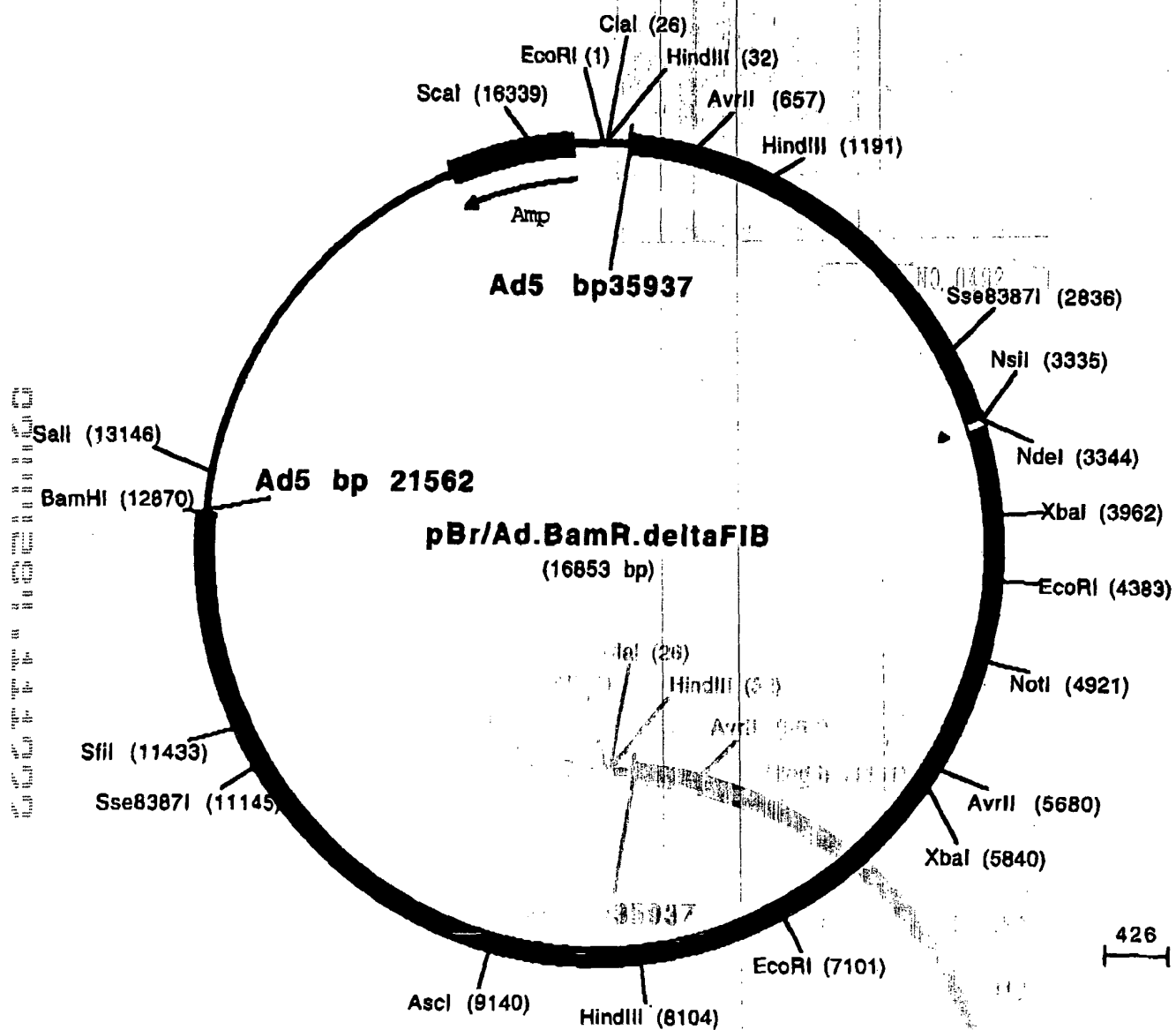


Figure 3

[illegible]

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 TAACAGACGAGCTGCTGCTGCTGCTGCTGCTGCTG
 TCAAACTAACTGCTGCTGCTGCTGCTGCTGCTG
 TGACTTAAGGCTGCTGCTGCTGCTGCTGCTGCTG
 ACCIACCTAACTGCTGCTGCTGCTGCTGCTGCTG
 CTAGAACCAAGCTGCTGCTGCTGCTGCTGCTGCTG
 ACTGCCCAAGCTGCTGCTGCTGCTGCTGCTGCTG
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[illegible][illegible]

Ad5/fib16 chimeric fiber

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TTCTGGAATGGCCTATGCTATGAATTTTTTCATGGTCTCTAAATGCAGAGGAAGCCCCGAAACTACC
GAAGTCACTCTCATTACCTCCCCCTCTTTTTTTCTTATATCAGAGAAGATGACTGAATGCATTAG

himeric f bar

[illegible]

Figure 4D: Sequence of Ad5/fib28 chimeric fiber

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chimeric fiber

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Figure 4E: Sequence

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CGTGCAAAACAATTCTCTCTCCTTAGGGGTTAACCCGCTTTTCTCATCACTGACTCTGGATTAGCT
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CGTGT'TTGACAACGAAGGGATACTAGCAAACAGTGCCACATGGGGTTATCGACAAGGACAGTCTGCC
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CTGAAGTTCAGAACATGGCTCTTACCTACACTTTTTTTGCAAGGTGACCCCTAACATGGCCATATCTTT
TCAGAGCATT'TATAATCATGCAATAGAAGGCTACTCTATTAAATTCNCCTGGCGCGTTCGAAATAAT
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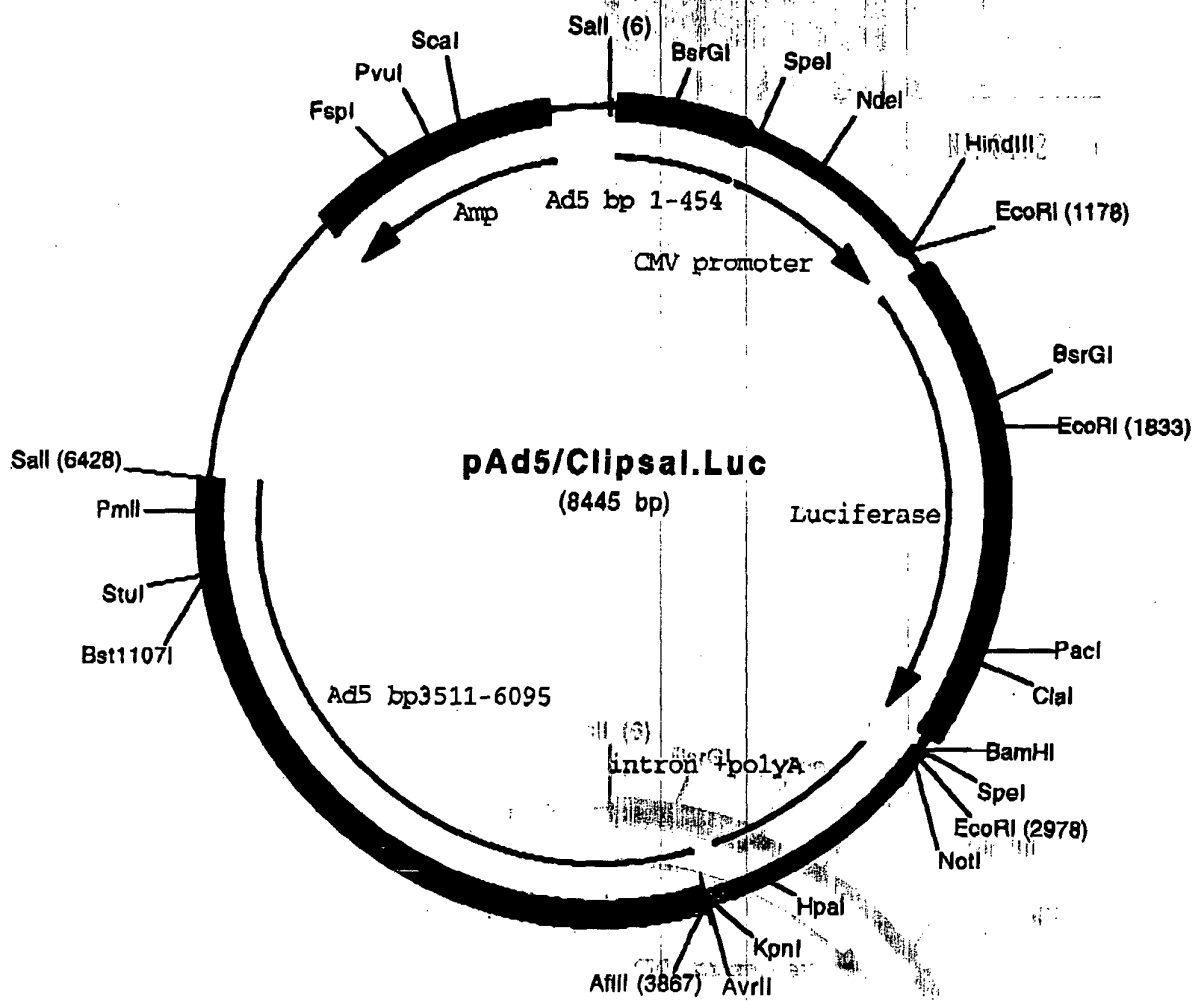
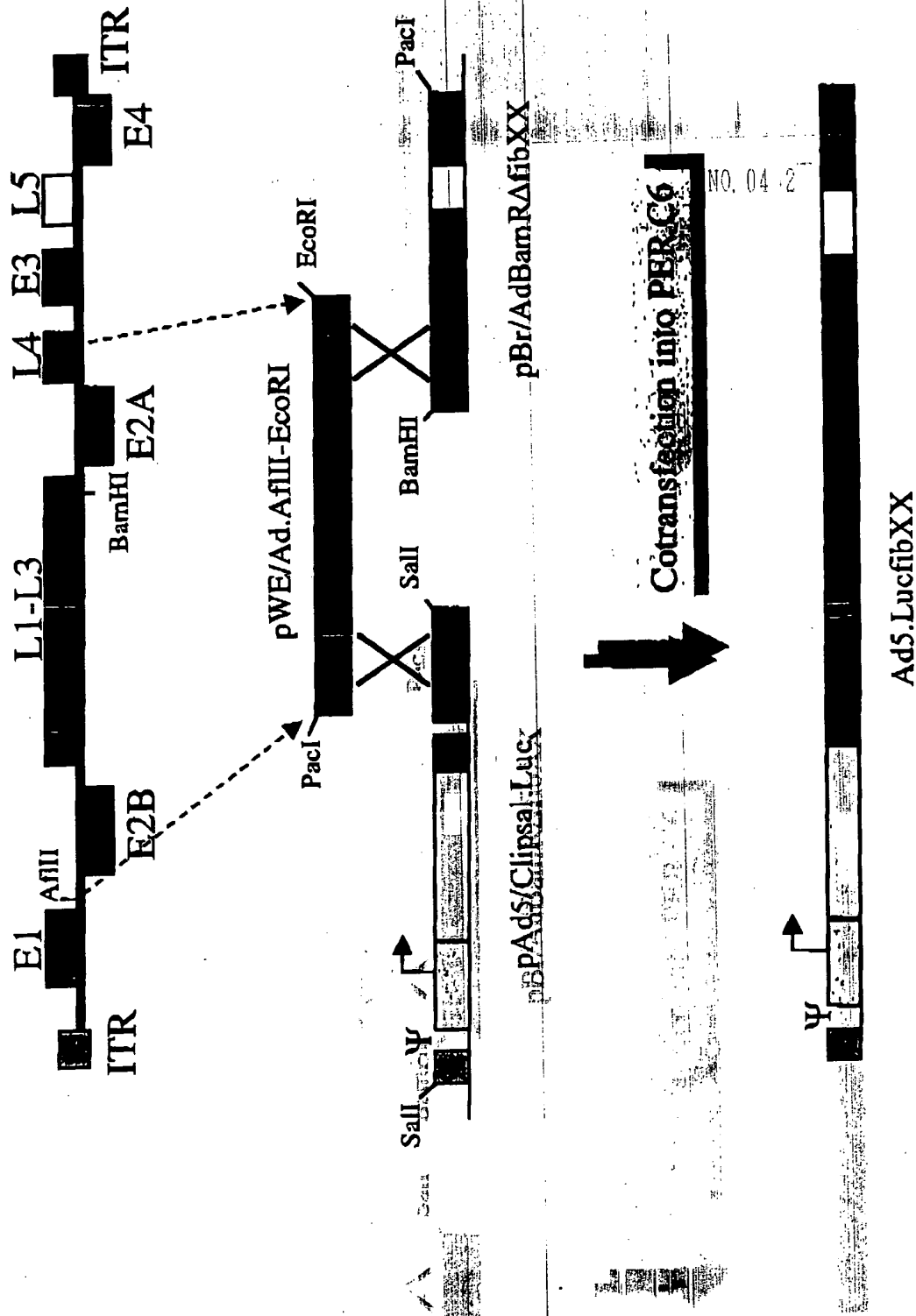


Figure 5

Figure 6: Generation of (chimeric) adenoviruses



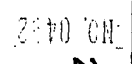
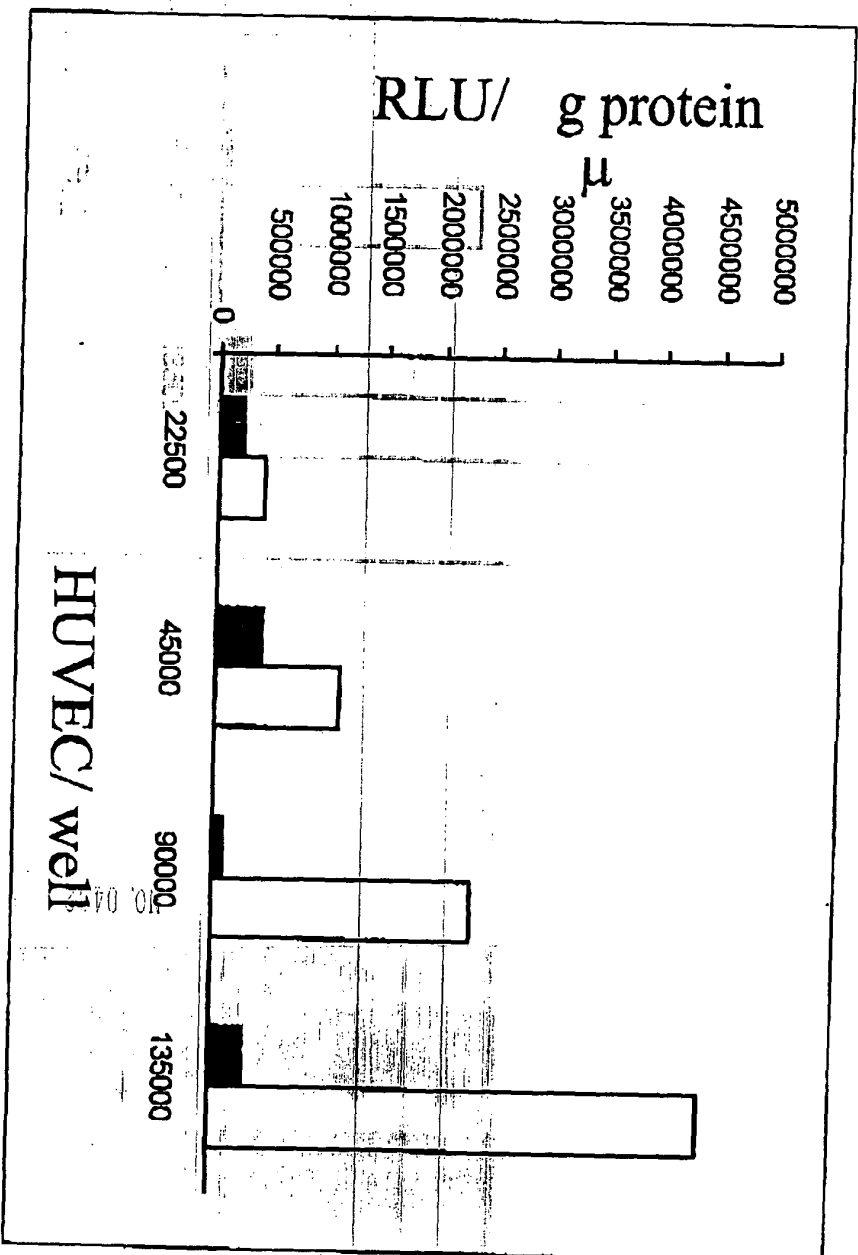
[illegible]

Figure 7b



Cellular uptake of protein

Figure 7c

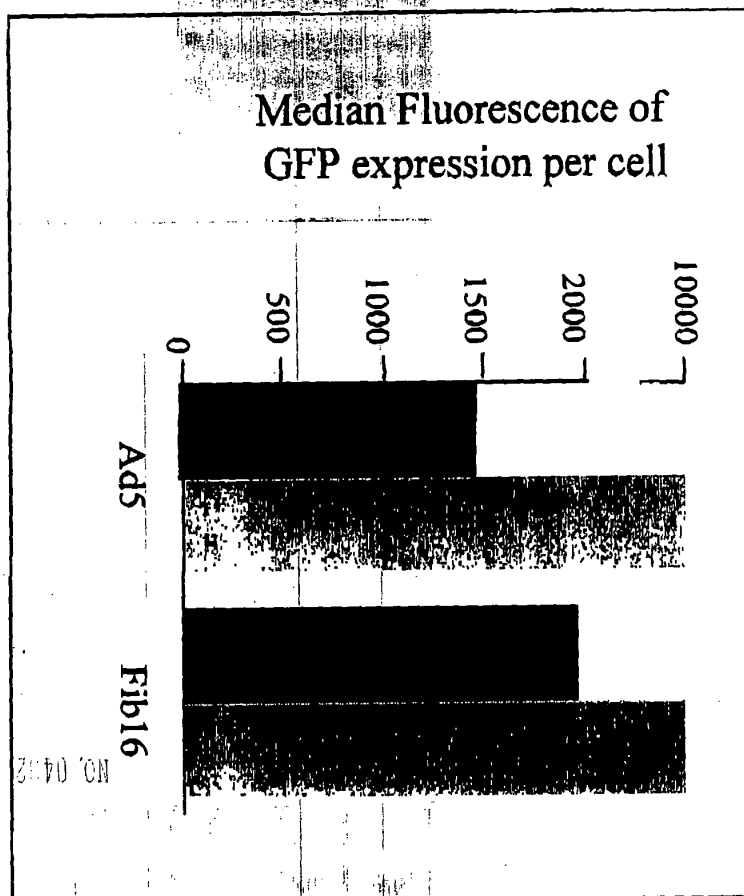
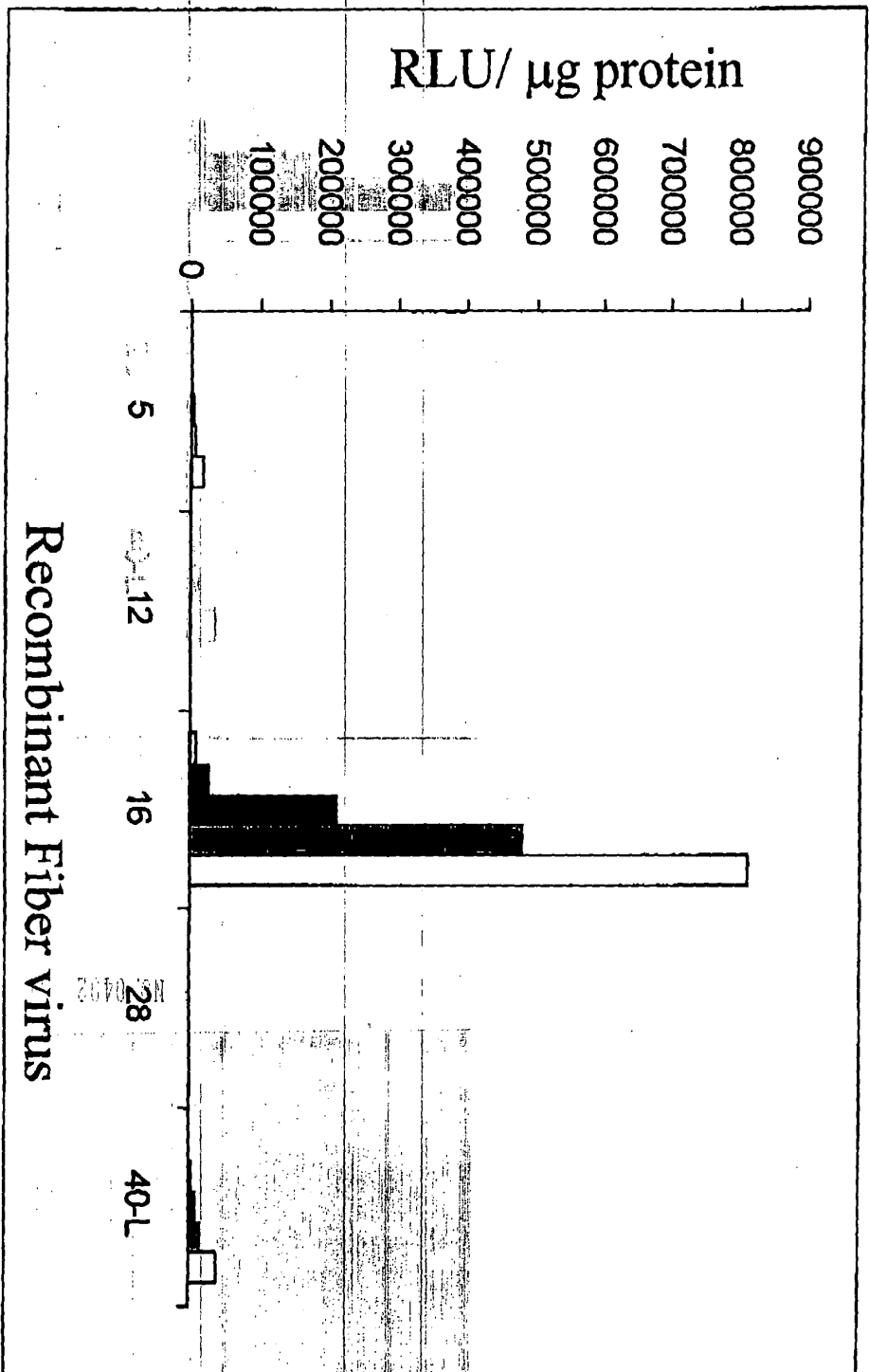


Figure 8a

[illegible]

RLU/ μg protein

HU Vsmc / well

HU Vsmc / well	RLU/ μg protein
0	~100,000
10,000	~2,800,000
20,000	~2,500,000
40,000	~2,800,000
60,000	~2,800,000
80,000	~2,800,000

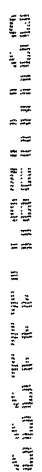


Figure 8c

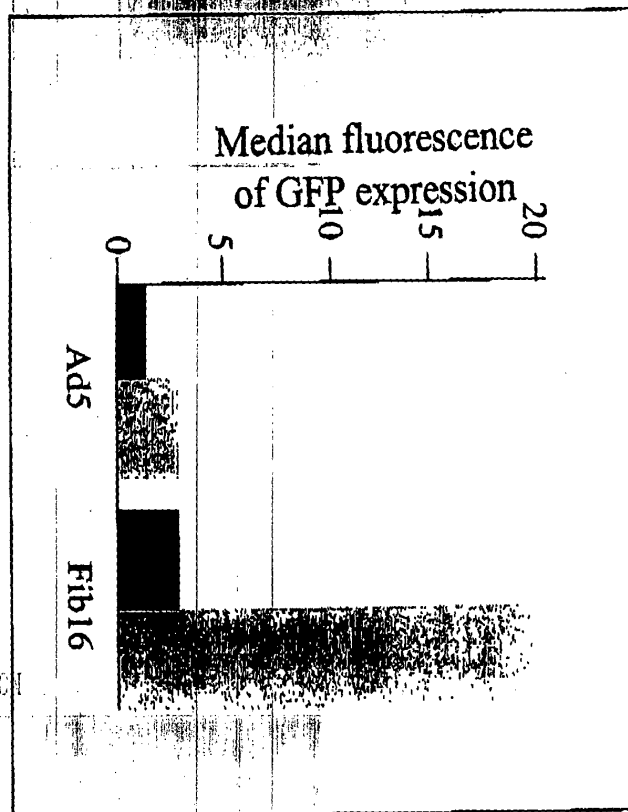
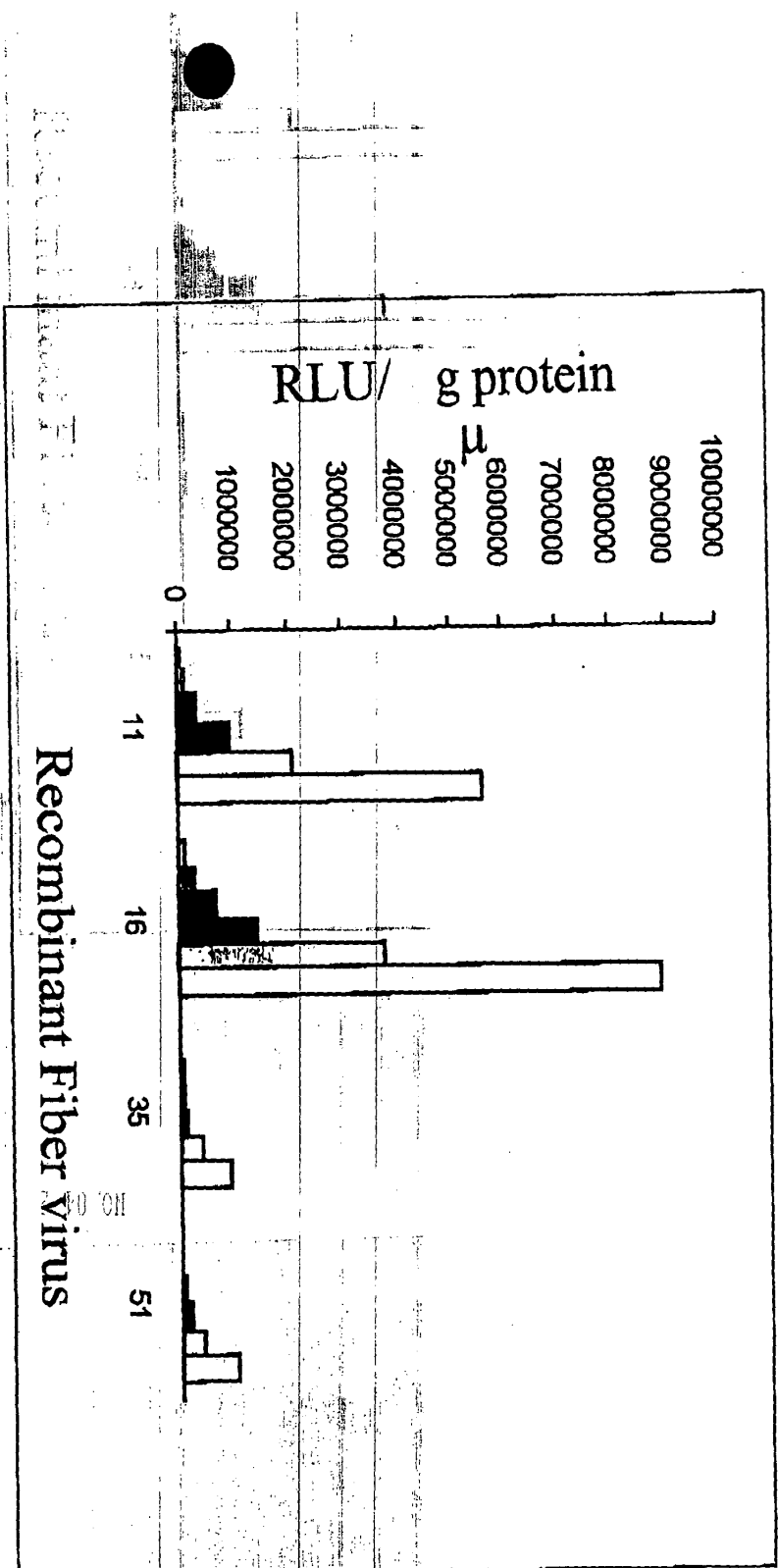
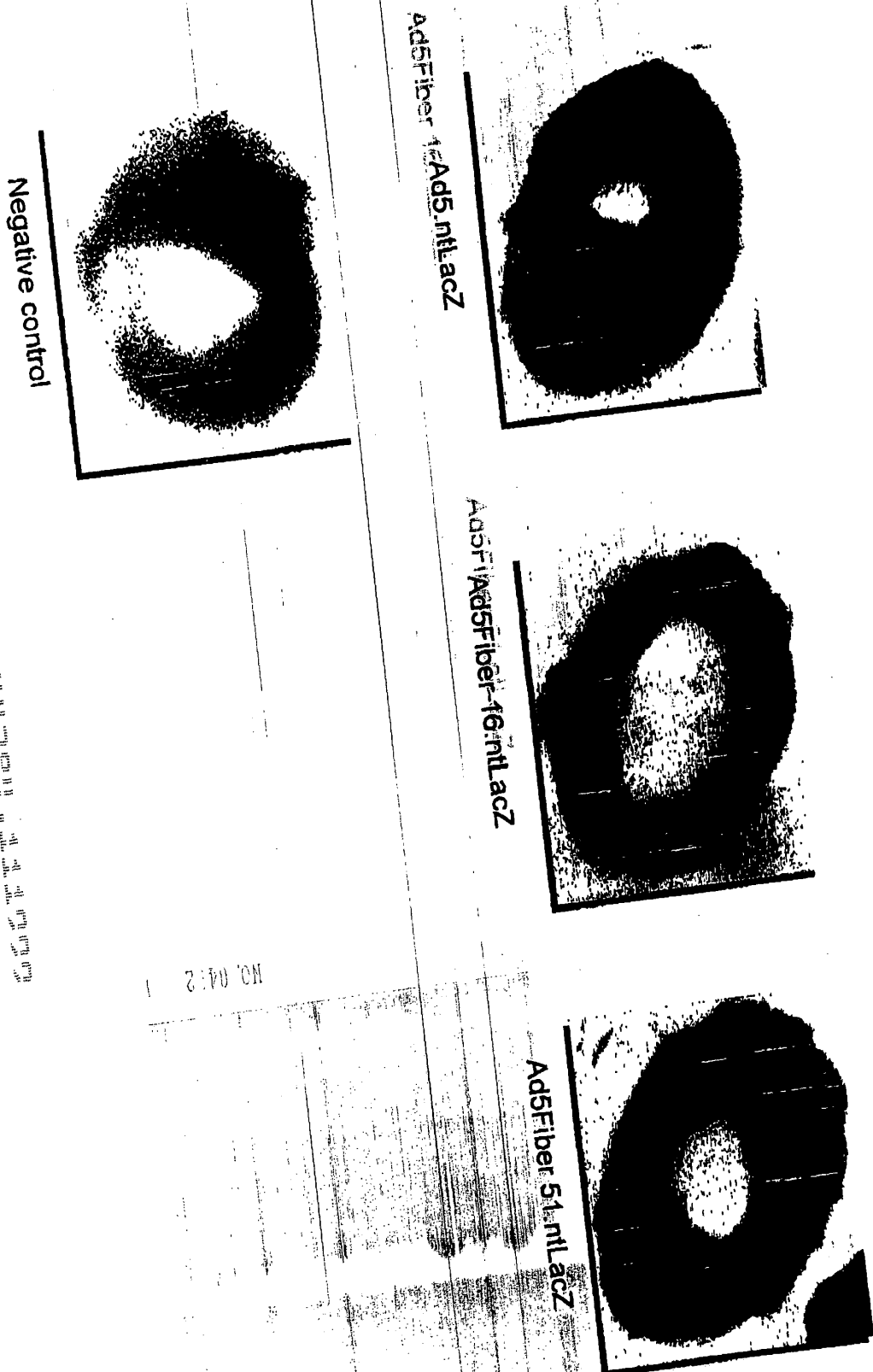


Figure 8d



100 80 60 40 20 0

Figure 8e:



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NO. 04:2

Ad5Fiber 51 nt

Ad5Fiber 51.ntlacZ

$\frac{1}{\sqrt{2}} \begin{pmatrix} 1 & i \\ 0 & 1 \end{pmatrix}$

REF ID: A6412

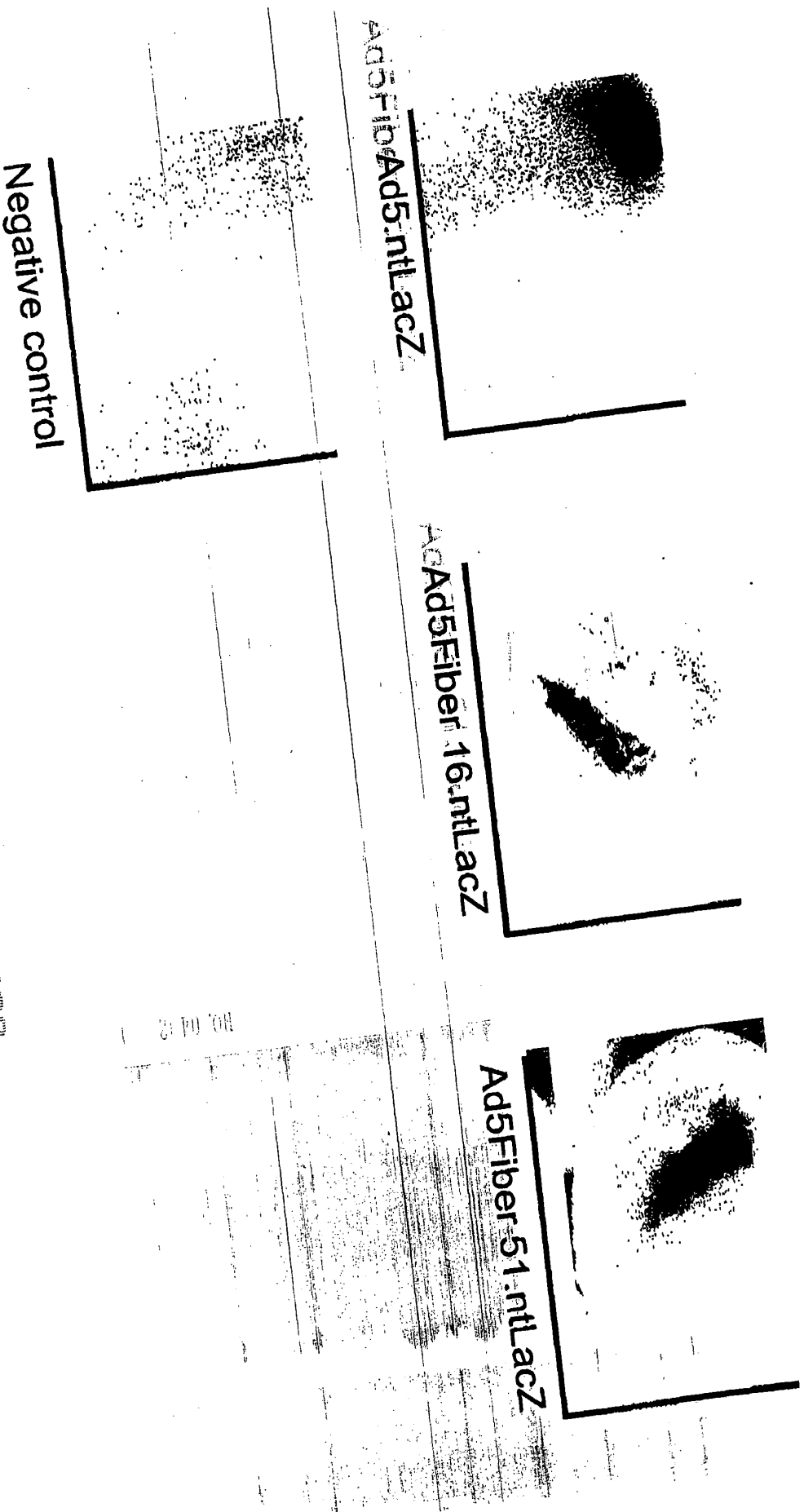
A vertical strip of three black and white photographs. The top photograph is a close-up of a person's face, showing their eyes and nose. The middle photograph is a close-up of a person's mouth, showing their lips and teeth. The bottom photograph is a close-up of a person's mouth, showing their lips and teeth.

Ad5Fiber Ad5.ntlacZ

Ad5Fiber 16.ntlacZ

[illegible]

Figure 8h



Thursday, November 10, 1994

1 ATGGC---CAAACGAGCTCGGCTAAGCAGACT--- Ad16 genbank.seq
1 ATGTTTGTTCAGATGAAGCGCGCAAGACCGTCTCTGAAGATA Ad5/fib16.seq

29 CCTTCAATCCGGTCTACCCCTATGAAGATGAAAGCAGCTC Ad16 genbank.seq
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121 GGTTTTTGACACAAAGCCCAGATGGAGTTCTAACTCTTAAAT Ad5/fib16.seq

149 GTGTTAATCCACTCACTACC GCCAGCGGACCCCTCCA ACT Ad16 genbank.seq
161 GTGTTAATCCACTCACTACC GCCAGCGGACCCCTCCA ACT Ad5/fib16.seq

189 TAAAGTTGG AAGCAGTCTTACAGTAGATACTATCGATGGG Ad16 genbank.seq
201 TAAAGTTGG AAGCAGTCTTACAGTAGATACTATCGATGGG Ad5/fib16.seq

229 TCTTTTGGAGGAAAATATAACTGCCGCAGCGCCACTCACTA Ad16 genbank.seq
241 TCTTTTGGAGGAAAATATAACTGCCGCAGCGCCACTCACTA Ad5/fib16.seq

269 AAAC TAACCACTCCATAGGTTTATTAAATAGGATCTGGCTT Ad16 genbank.seq
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309 GCAAACAAGGATGATAAACTTTGTTTATCGCTGGGAGAT Ad16 genbank.seq
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Fig 9A.

Alignment Report of Untitled, using Clustal method with Weighted residue weight table.
Thursday, November 19, 1998 18:26

Page 2

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869  ACTACAAATCTACCAATGGAACCTCTCTTTCCACTAAAAGT Ad16 genbank.seq
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989  AAGCCCCGGAAACTACCGAAGTCACTCTCATTACCTCCCC Ad16 genbank.seq
1001 AAGCCCCGGAAACTACCGAAGTCACTCTCATTACCTCCCC Ad5/fib16.seq

1029 CTTCTTTTTTTTCTTATATCAGAGAAAGATGACTGA Ad16 genbank.seq
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Decoration 'Decoration #1': Box residues that differ from Ad16 genbank.seq

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CCATTTTCTACCTGCGCC
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ACATTTTCTACCTGCGCC
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TTCTTTTCTACCTGCGCC
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end of Ad16

Page :

Decoration 'Decoration #1': Box residues that differ from the Consensus.

[illegible]